

- [0253] 23. Shahi, P., Kim, S. C., Haliburton, J. R., Gartner, Z. J. & Abate, A. R. Abseq: Ultrahigh-throughput single cell protein profiling with droplet microfluidic barcoding. *Scientific Reports* 2016 6 7, 44447 (2017).
- [0254] 24. Butler, A., Hoffman, P., Smibert, P., Papalexi, E. & Satija, R. Integrating single-cell transcriptomic data across different conditions, technologies, and species. *Nature Biotechnology* 201129:1136, 411-420 (2018).
- [0255] 25. Langfelder, P. & Horvath, S. WGCNA: an R package for weighted correlation network analysis. *BMC Bioinformatics* 9, 1 (2008).
- [0256] 26. Thomas, G. D. et al. Deleting an Nr4a1 Super-Enhancer Subdomain Ablates Ly6C^{low} Monocytes while Preserving Macrophage Gene Function. *Immunity* 45, 975-987 (2016).
- [0257] 27. Barry, K. C. et al. A natural killer-dendritic cell axis defines checkpoint therapy-responsive tumor microenvironments. *Nature Medicine* 2018 24, 1178-1191 (2018).
- [0258] 28. Singhal, S. et al. Human tumor-associated monocytes/macrophages and their regulation of T cell responses in early-stage lung cancer. *Science Translational Medicine* 11, eaat1500 (2019).
- [0259] 29. Zhu, Y. P. et al. Identification of an Early Unipotent Neutrophil Progenitor with Pro-tumoral Activity in Mouse and Human Bone Marrow. *Cell Reports* 24, 2329-2341.e8 (2018).
- [0260] 30. Li, H. Y. et al. The Tumor Microenvironment Regulates Sensitivity of Murine Lung Tumors to PD-1/PD-L1 Antibody Blockade. *Cancer Immunol Res* 5, 767-777 (2017).
- [0261] 31. Broz, M. L. et al. Dissecting the Tumor Myeloid Compartment Reveals Rare Activating Antigen-Presenting Cells Critical for T Cell Immunity. *Cancer Cell* 26, 638-652 (2014).
- [0262] 32. Olingy, C. E. et al. Non-classical monocytes are biased progenitors of wound healing macrophages during soft tissue injury. *Scientific Reports* 2016 6 7, 447 (2017).
- [0263] 33. Loyher, P.-L. et al. Macrophages of distinct origins contribute to tumor development in the lung. *Journal of Experimental Medicine* 20180534 (2018).
- [0264] 34. Blackburn, S. D. et al. Coregulation of CD8⁺ T cell exhaustion by multiple inhibitory receptors during chronic viral infection. *Nature Immunology* 201112:8 10, 29-37 (2009).
- [0265] 35. Green, D. S. et al. A Phase 1 trial of autologous monocytes stimulated ex vivo with Sylatron® (Peginterferon alfa-2b) and Actimmune® (Interferon gamma-1b) for intra-peritoneal administration in recurrent ovarian cancer. *Journal of Translational Medicine* 2018 16:116, 196 (2018).
- What is claimed:
1. A method of determining whether a subject will respond to a treatment for a neoplasia, neoplastic disorder, tumor, cancer or malignancy, the method comprising detecting the amount of CD33 in a biological sample from the subject, comparing the measured amount to a reference amount, wherein a modified measured amount compared to the reference amount is indicative that the subject will or will not respond to the treatment for a neoplasia, neoplastic disorder, tumor, cancer or malignancy.
 2. The method of claim 1, wherein the modified measured amount is an increase in the measured amount.
 3. The method of claim 1, wherein the modified measured amount is a decrease in the measured amount.
 4. The method of claim 2, wherein the increase in the measured amount is indicative that the subject will respond to the treatment for a neoplasia, neoplastic disorder, tumor, cancer or malignancy.
 5. The method of claim 3, wherein the decrease in the measured amount compared to the reference amount is indicative that the subject will not respond to the treatment for a neoplasia, neoplastic disorder, tumor, cancer or malignancy.
 6. The method of claim 2 or 4, wherein the measured amount is CD33 high expression in the biological sample.
 7. The method of claim 3 or 5, wherein the measured amount is CD33 low expression in the biological sample.
 8. The method of claim 2 or 4, wherein the measured amount is CD33 low expression in the biological sample.
 9. The method of claim 3 or 5, wherein the measured amount is CD33 high expression in the biological sample.
 10. A method of determining whether a subject will respond to a treatment for a neoplasia, neoplastic disorder, tumor, cancer or malignancy, the method comprising detecting the presence or amount of CD33 expressing cells in a biological sample from the subject, comparing the measured presence or amount to a reference presence or amount, wherein a modified measured presence or amount as compared to the reference presence or amount is indicative that the subject will or will not respond to the treatment for a neoplasia, neoplastic disorder, tumor, cancer or malignancy.
 11. The method of claim 10, wherein the CD33 expressing cells measured are cells with high CD33 expression.
 12. The method of claim 10, wherein the CD33 expressing cells measured are cells with low CD33 expression.
 13. The method of claim 11, wherein the measured presence or amount of cells with high CD33 expression is indicative that the subject will respond to the treatment for a neoplasia, neoplastic disorder, tumor, cancer or malignancy.
 14. The method of claim 12, wherein the measured presence or amount of cells with low CD33 expression is indicative that the subject will not respond to the treatment for a neoplasia, neoplastic disorder, tumor, cancer or malignancy.
 15. A method of determining whether a subject will respond to a treatment for a neoplasia, neoplastic disorder, tumor, cancer or malignancy, the method comprising detecting the presence or measuring the amount of cells with high expression of CD33 in a biological sample from the subject, wherein the presence of cells with high expression of CD33 or a high amount of cells with high expression of CD33 indicates that the subject will respond to the treatment for a neoplasia, neoplastic disorder, tumor, cancer or malignancy.
 16. The method of claim 15, wherein the method comprises detecting the presence of or measuring the amount of CD33^{hi} myeloid cells in the biological sample.
 17. The method of any preceding claim, wherein the method comprises detecting the presence or measuring the amount of classical or non-classical monocytes in the biological sample.
 18. The method of claim 17, wherein the method comprises detecting the presence or measuring the amount of classical monocytes in the biological sample.